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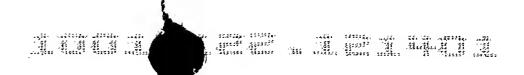
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Arg Phe Ser Leu Val Ser Gly Trp Gly Gln Leu Leu Asp Arg Gly Ala 340 345 350 Thr Ala Leu Glu Leu Met Val Leu Asn Val Pro Arg Leu Met Thr Gln 355 365 360 Asp Cys Leu Gln Gln Ser Arg Lys Val Gly Asp Ser Pro Asn Ile Thr 370 375 380 Glu Tyr Met Phe Cys Ala Gly Tyr Ser Asp Gly Ser Lys Asp Ser Cys 385 390 395 400 Lys Gly Asp Ser Gly Gly Pro His Ala Thr His Tyr Arg Gly Thr Trp 405 410 415 Tyr Leu Thr Gly Ile Val Ser Trp Gly Gln Gly Cys Ala Thr Val Gly 420 425 430 His Phe Gly Val Tyr Thr Arg Val Ser Gln Tyr Ile Glu Trp Leu Gln 435 440 445 Lys Leu Met Arg Ser Glu Pro Arg Pro Gly Val Leu Leu Arg Ala Pro 450 455 460 Phe Pro 465 <210> 3 <211> 31 <212> DNA <213> Homo sapiens <400> 3 gctgcaggtg cgtccaggga ggttttctcc a 31 <210> 4 <211> 31 <212> DNA <213> Homo sapiens <400> 4 ctcctgtcgg tgccatgagg ggtactctct g 31